

PRINTER RUSH
(PTO ASSISTANCE)

Application : 09/744,167 Examiner : Borin GAU : 1631

From: J. Robbins Location: (IDC) FMF FDC Date: 1-11-06

Tracking #: 06041103 Week Date: 11-15-04

DOC CODE	DOC DATE	MISCELLANEOUS
<input type="checkbox"/> 1449	_____	<input type="checkbox"/> Continuing Data
<input type="checkbox"/> IDS	_____	<input type="checkbox"/> Foreign Priority
<input type="checkbox"/> CLM	_____	<input type="checkbox"/> Document Legibility
<input type="checkbox"/> IIFW	_____	<input type="checkbox"/> Fees
<input type="checkbox"/> SRFW	_____	<input checked="" type="checkbox"/> Other <u>Let. 11-3-2005</u>
<input type="checkbox"/> DRW	_____	
<input type="checkbox"/> OATH	_____	
<input type="checkbox"/> 312	_____	
<input type="checkbox"/> SPEC	_____	

[RUSH] MESSAGE: Page 62, Tables 9 and 10 which
were provided per previous response to XRUSH. (see
Let. 11-3-2005) are still illegible for publication

[XRUSH] RESPONSE: _____

Dch

clearest copy provided

INITIALS: [Signature]

NOTE: This form will be included as part of the official USPTO record, with the Response document coded as XRUSH.

Table 9

hSARA	MFNDENVAEDQLIKKMYSDDCSAVEGERKCGMADLPDEKNVLVVAVHHNCDKRTQNDLQDNNYNSOSLND	77
XSARA	MPKVVGGDTMAEDSLFNNIGPSEIVNPSTVSS-----SLQALDDQS-----VNIHNEKSVILLADGFSPSSP-----KSIIN	70
hSARA	AFSLDNEENRQTDQFSFSINESTKDNINSEKNDPLNRPKTEGRSVNHCPTSSPSLASVCSFSQLDDGSIERDPSMS	157
XSARA	FLCLTHDN-----EHLHSONSVDDNDK-----ETVLTISVLPTIIQDTSNVSEDPAINKP-----GPKEPHR	127
hSARA	ATTSITVDSVISSCGTDGFAVKKDEN-YLPDELTGKISSPRFDLGEPSNFSHASEGILMKKEPAEESTTEESIRSGLP	236
XSARA	ALKETTSVILPEIKPYSTCAALSFERANKVPSYQLN-----NTDILS-----VSPVVEACSEQQQKHTSSLHEEKLEFGVS	198
hSARA	LLLKPDMPNGSGRNNDCERCSDCLVPNEVRADENEGYEHEETLCTTEFLNNTTEFSESQDNTMVKLTKLNEFNDSQVNEE	316
XSARA	ATES-----FAATAETVLONEALSAEFTDIUVKNFSDSCVINGDLTKSCGLS-----DES	250
hSARA	KEKFLQISPEEDTNGDSGGQCVGLADAGLDDGTCTISESECDPSTVIDTPRANVLSNGCDSYGMQDPCGVSEVPKTLPSK	396
XSARA	NEKFCASKETFE-----CG-----VDANVILENACVAYKKAIDLPEENGQINAPMELLYNGCDSYGMKNFAVAQNPKNLPSK	319
hSARA	EDSVTEEKEIEESKSEYVNIYEQ-RGNEATESSGLLNNSTCDLHKKNYLMHFCSQVPSVLGQSSPKVVASLPSISVFPFG	475
XSARA	EDSVTEEKEIEESKSEYVTGVYEQPEDVTERGGLLLNAKADQMKNN-LHSLOQVPSMAGQTSPPKGGKIVQSLSVPYG	398
hSARA	GARPQPSNLKLQIPKPLSDHLND-FFANSGNNTTKKNDILGKAKLGENSATNVCSPLGNISNVDTNGEHLESYEAE	553
XSARA	GARTKQPTHKLHLPKPLSEMLQSLDIPNACSSKYKNDMLNKSNOGDMLISESLREDSAVRSPTVDANGDFPGEYRGP	478
hSARA	ISTRPLALAPDSPDNOLRAGQFGISARKPFTTLGEVAPVWVPDSQAPNCMKCEARFTFTKRHHCRACGKVFCASCCSL	633
XSARA	GS--LCLAVSPDSPDNOLRAGQFGVVISKPFPTTLGEVAPVWVPDSQAPNCMKCEARFTFTKRHHCRACGKVFCACCSL	556
hSARA	KCKILYMDRKEARVCVICHSVLMNAQAWENMHSSASSPNNPNAEYCSPTIPPLQQAQASGALSSPPPTVNVVGVGLKHP	713
XSARA	KCKILOYMDKKEARVCVICHSVLMNAQAWENMLSAVSQSPNNPNAEYCSPTIPPMQQAQASGALSSPPPTVNVVGVGLKHP	636
hSARA	GAEVACREQRRVWFADGILPNGEVADAAKITNGTSSAGTLAVSHDPVKPVTHISPLPAETDILFSGSITQVGSVPVGEA	793
XSARA	GTESSCKEORRVWFADGILPNGETADSD--NNNVTVAGTLTVSHNNNSTSSS-----ENTSGCGSITQVG-----SA	705
hSARA	MNLIPEDGLPILISTGVKGDYAVEEKPSQISVMHQLEDGGPDPLVFLNANLLSHVKIVNYVNRKWCFTTKGHHAVGQ	873
XSARA	MNLIPEDGLPILISTGVKGDYAVEERPQMSVMHQLEEGPDPLVFLNANLLAHVKIVNYVNRKWCFTTKGHHAVGQ	785
hSARA	SEIVILLQCLPDEKCLPKDIEHMFVQLYRDALAGNVVNLGHSFSSQSLGSKHEGGFLYVITYQSLQDLVLPTPEYLF	953
XSARA	AEIVILLQCLPDEKCLPRDLESIFVELYQEAIAAGNVVNLGHSFSSQSLGSKDHGGFLYVAITYQSLQDLVLPAPEYLF	865
hSARA	GILIQKWETPWAKVFFIRLNLRLGAEYRLYPCPLFSVRFRKPLPGETGHTIMNLLADFRNYQYTLPPVQGLVVDMEVRKT	1033
XSARA	GILIQKWETPWAKVFFIRLNLRLGAEYRLYPCPLFSVRYRKPLPGETGHTIINVLADFRNYQYTLPPVQGLVVDMEVRKT	945
hSARA	SIKIPSNRYNEMHKAMNKSNEHVLAAGACFNEKADSHLVCVQNDGNYQTQAISIHQPRKVTGASFFVFGALKSSSGY	1113
XSARA	SIKIPSNRYNEMHKAMNKSNEHVLAAGACFNOHADSHLVCVQNDGNYQTQAISIHQPRKVTGASFFVFGALKSSSGY	1025
hSARA	LAKSSIVEDGVHVQITAEIMDSLRQALREMKDFTITCGKADAEEDEHVIHQWDDDKNVSKGVSPIDGKSMETITNVK	1193
XSARA	LAKSSIVEDGVHVQITAEIMDALRQSLREMKDFTITCGKADAEEDEHVYVQWVEDDKNVKGVSPIDGKSMESVTSVK	1105
hSARA	IFHGSEYKANKSVIRWTEVFFLENDQDQNLSDPADHSRLTEHVAKAFCLALCPHLKLLKEDGMTRLGLRVTLDSQVGY	1273
XSARA	IFHGSEYKANKSVIRWTEVFFLDNEEQQSLSDPADHSRLTENVAKAFCLALCPHLKLLKEDGMTRLGLRVSLDSQVGY	1185
hSARA	QAGSNGQFLPSQYINDLDSALVPVTHGGACQLSEGPVVMELIFYILENTV- 1323	
XSARA	QAGSNGQFLPARYINDLDSALVPVTHGGTCQLSEGPVVMELIFYILENTS- 1235	

Table 10

hSARA	587	GEVAPVWVPDSQAPNCMKCEARFTFTKRHHHCRACGKVFCASCCSLKCKLYMDRKEARVCVICHSVI	655
XSARA	510	LGEVAPVWVPDSQAPNCMKCEARFTFTKRHHHCRACGKVFCACCSLKCKLYMDRKEARVCVICHSVI	578
KIAA0305	737	LGCKDPVWVPDSQAPNCMKCEARFTFTKRHHHCRACGKVFCACCSLKCKLYMDRKEARVCVICHSVI	800
FGD1	720	LGRAPTHIREKEVIMCMHCEPIFNSITKRHHHCRACGKVFCACCSLKCKLYMDRKEARVCVICHSVI	485
Hrs	153	AAERAPVWD--AEECHRCVQF-GVTRKHHHCRACGQIFCGKCSSKYSTIPKFGIEKEVRVCEPCYEOL	219
Hrs-2	153	AAERAPVWD--AEECHRCVQF-GVTRKHHHCRACGQIFCGKCSSKYSTIPKFGIEKEVRVCEPCYEOL	219
EEA-1	1341	TOALNRKMAEDNEVQNCMACGKEFSVTVRRHHCRACGQIFCAECSAKNALTPSSKILPVAVCDACFNDL	1408

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